



The first conservation geneticist

# THE VARIATION OF ANIMALS AND PLANTS UNDER DOMESTICATION

BY
CHARLES DARWIN, M. A., F. R. S., ETC.

### CHAPTER XVII.

ON THE GOOD EFFECTS OF CEOSSING, AND ON THE EVIL EFFECTS OF CLOSE INTERBREEDING.

DEFINITION OF CLOSE INTERBREEDING—AUGMENTATION OF MOBBID TENDENCIES—GENERAL EVIDENCE OF THE GOOD EFFECTS DERIVED FROM CROSSING, AND ON THE EVIL EFFECTS FROM CLOSE INTERBREEDING—ACTILE, CLOSELY INTERBREED HALF-WILD CATTLE LONG KEFT IN THE SAME PARKS—SHEET—FALLOW-DEER—DOGS, RABBITS, PIGS—MAN, ORIGIN OF HIS ABBORNENCE OF INCESTUOUS MARRIAGES—FOWLS—FIGEONS—HIVE-BEES—VLANTS, GENERAL CONSIDERATIONS ON THE ENSETTS DERIVED FROM CROSSING—MELONS, FRUITTREES, PLS, CABBAGES, WHEAT. AND FORESTTREES—ON THE INCREASED SIZE OF HYBRID PLANTS, NOT EXCLUSIVELY DUE TO THEIR STERILITE—ON CRETAIN PLANTS WHICH EITHER NOR-MALLY OR ABNORMALLY ARE SELF-IMPOTENT, BUT ARE FEITILE, BOTH ON THE MALE AND FEMALE SIDE, WHEN GROSSED WITH DISTINCT INDIVIDUALS EITHER OF THE SAME OR ANOTHER SPECIES—CONCLUSION.

That any evil directly follows from the closest interbreeding has been denied by many persons; but rarely by any practical breeder; and never, as far as I know, by one who has largely bred animals which propagate their kind quickly. Many physiologists attribute the evil exclusively to the combination and consequent increase of morbid tendencies common to both parents; and that this is an active source of mischief there can be no doubt.

Charles Darwin (1898)

As some of our British parks are ancient, it occurred to me that there must have been long continued close interbreeding with the fallow deer (*Cervus dama*) kept in them; but on inquiry I find that it is a common practice to infuse new blood by procuring bucks from other parks.

Charles Darwin (1896, page 99)



### **CONSERVATION GENETICS**

- (1) Management and reintroduction of captive populations, and the restoration of biological communities.
- (2) Description and identification of individuals, genetic population structure, kin relationships, and taxonomic relationships.
- (3) Detection and prediction of the effects of habitat loss, fragmentation, and isolation.
- (4) Detection and prediction of the effects of hybridization and introgression.
- (5) Understanding the relationships between adaptation or fitness and genetic characters of individuals or populations.
- (6) Application of genetic markers to understand the basic biology of species and populations (including invasive species).
- (7) Forensics.
- (8) Application of genetic engineering to problems in conservation.

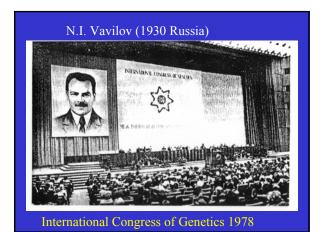
# **History of Conservation Genetics**

N.I. Vavilov (1930 Russia)
Plant collections

Paavo Voipio (1950 Finland) Game management

Sir Otto Frankel (1970 Australia)
Preservation of agricultural diversity
Conservation of natural populations

Michael Soulé (1980 USA)
Founder of conservation biology



PAPERS ON GAME RESEARCH

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Evolution at the population level with special reference to game animals and practical game management

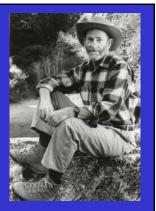
by

Paavo Yoipio.

To be presented, with the permission of the Section of Mothematics and Neutral Sciences of the Philosophical Foculty of the University of Minimals, for public criticism in Auditorium 12, October 213¢ 1950, or 12 o'clock.

Symposium on Genetics and Society: XIII International Congress of Genetics GENETIC CONSERVATION: OUR EVOLUTIONARY RESPONSIBILITY O. H. FRANKEL CSIRO, Division of Plant Industry, P.O. Box 1600, Canberra City. A.C.T. Australia The time scale of concern Operator Time scale Wild-life to 8,000 BC hunter-gatherer next meal 1 day Domesticated plants to 1850 AD "primitive" or "traditional" the next crop 1 year peasant farmer from 1850 plant breeder the next variety 10 years from 1900 crop evolutionist to broaden the 100 years genetic base Wildlife genetical conservationist dynamic wildlife 10,000 years + current public interest politician next election

Soulé, M. E., and Wilcox, B. M. (Editors). 1980. Conservation Biology An Evolutionary -Ecological Perspective. Sinauer, Sunderland, MA.



# Participant Background Survey

There are many different molecular techniques (allozymes, mtDNA, microsatellites). What are the best methods for different questions?

How can results of genetic studies be misinterpreted?

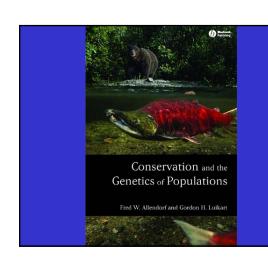
Is a difference in chromosome number sufficient to make decisions in taxonomy at the species, subspecies, or population level?

When should groups be managed as separate populations versus increasing genetic exchange to avoid inbreeding depression?

How large does a population have to be to avoid detrimental genetic effects?

I am interested in developing a better understanding of evolutionarily significant units and distinct populations segments.

Understanding concepts of adaptation and evolution in species of invasive plants.



# Conservation Genetics Joke of the Day

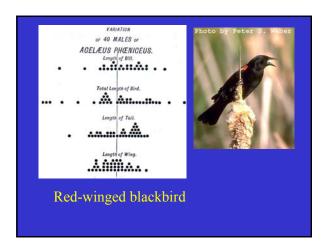


## PHENOTYPIC VARIATION IN NATURAL POPULATIONS



Few persons consider how largely and universally all animals are varying. We know however, that in every generation, if we would examine all the individuals of any common species, we should find considerable differences, not only in size and color, but in the form and proportions of all the parts and organs of the body.

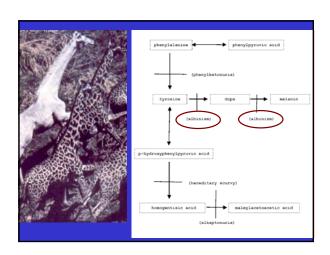
Alfred Russel Wallace



The many slight differences which appear in the offspring from the same parents, or which it may be presumed have thus arisen, from being observed in the individuals of the same species inhabiting the same confined locality, may be called individual differences. These individual differences are of the highest importance for us, for they are often inherited, as must be familiar to every one; and they thus afford materials for natural selection to act on and accumulate, in the same manner as man accumulates in any given direction individual differences in his domesticated productions. I am convinced that the most experienced naturalist would be surprised at the number of the cases of variability, even in important parts of structure, which he could collect on good authority, as I have collected, during a course of years. It should be remembered that systematists are far from being pleased at finding variability in important characters, and that there are not many men who will laboriously examine internal and important organs, and compare them in many specimens of the same species.

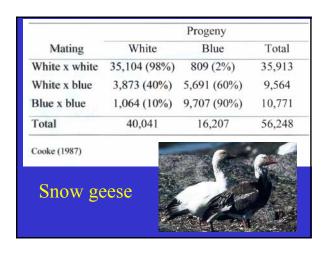


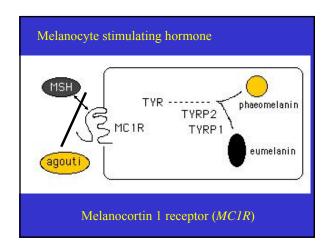


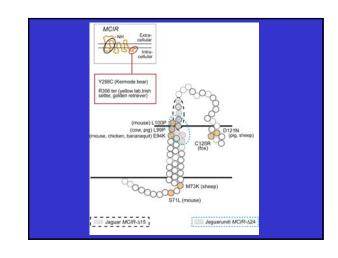


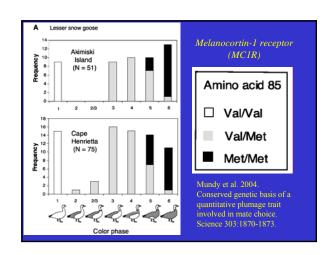


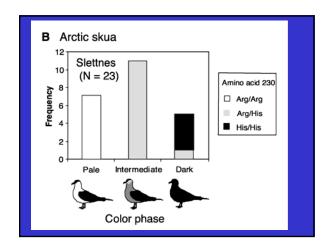
	Number	Pro	geny	
Mating	of families	Red	Gray	
Red x red	8	23	5	
Red x gray	46	68	63	
Gray x gray	135	0	439	
200	學			





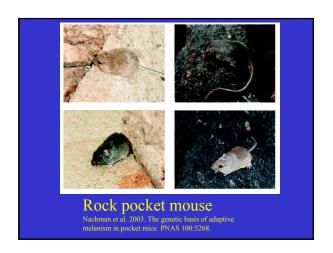


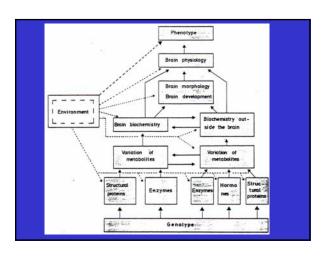


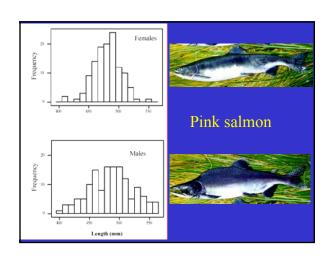












$$V_P = V_G + V_E$$
  
Heritability =  $V_G / V_P$ 

